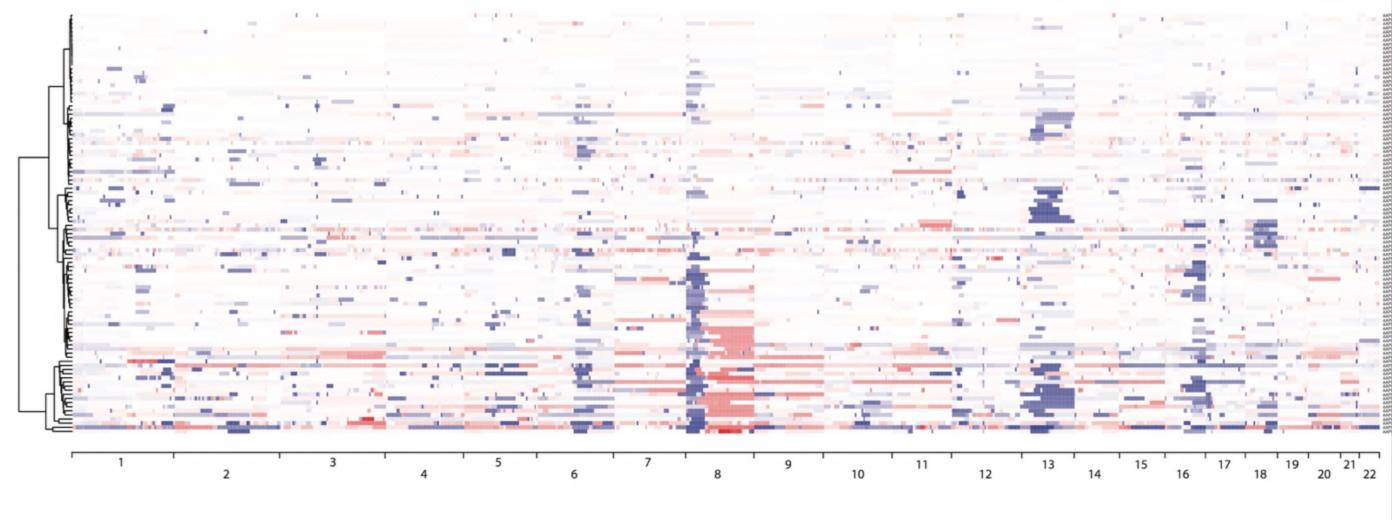


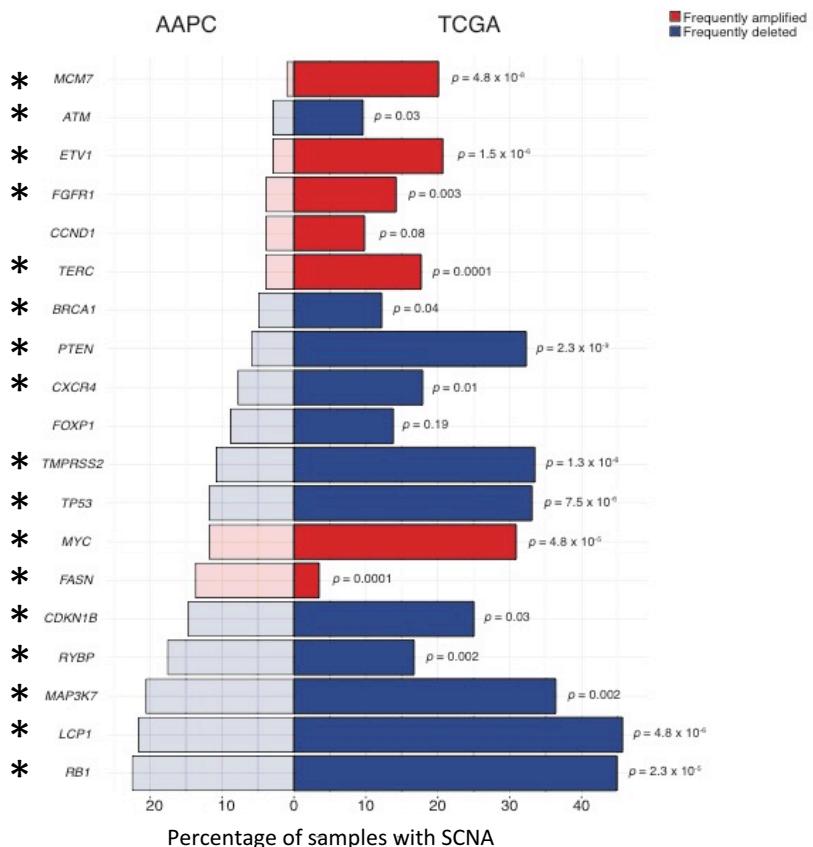
Supplemental Figure 1

log2
copy
ratio
3
2
1
0
-1

A.

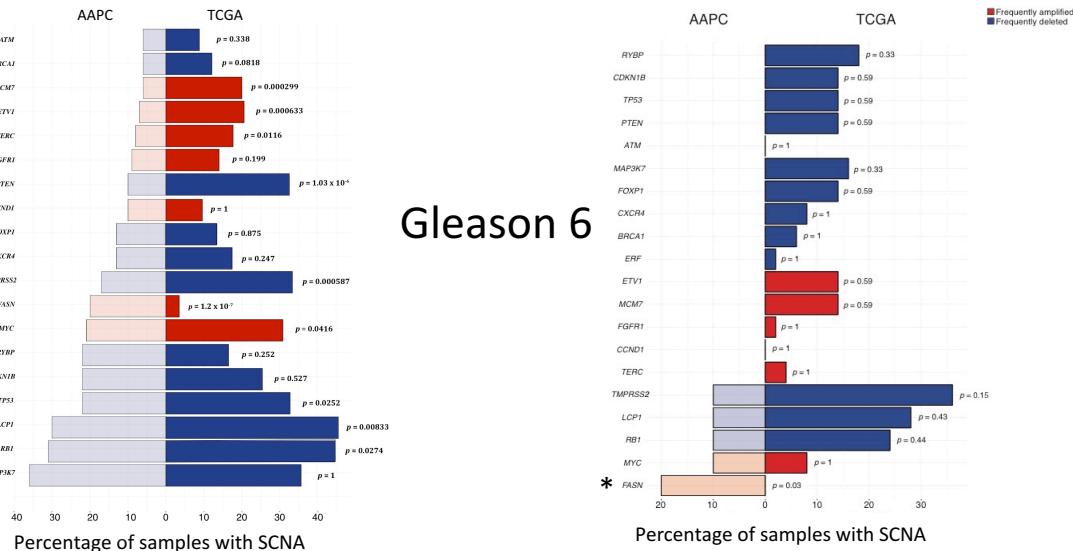


B.

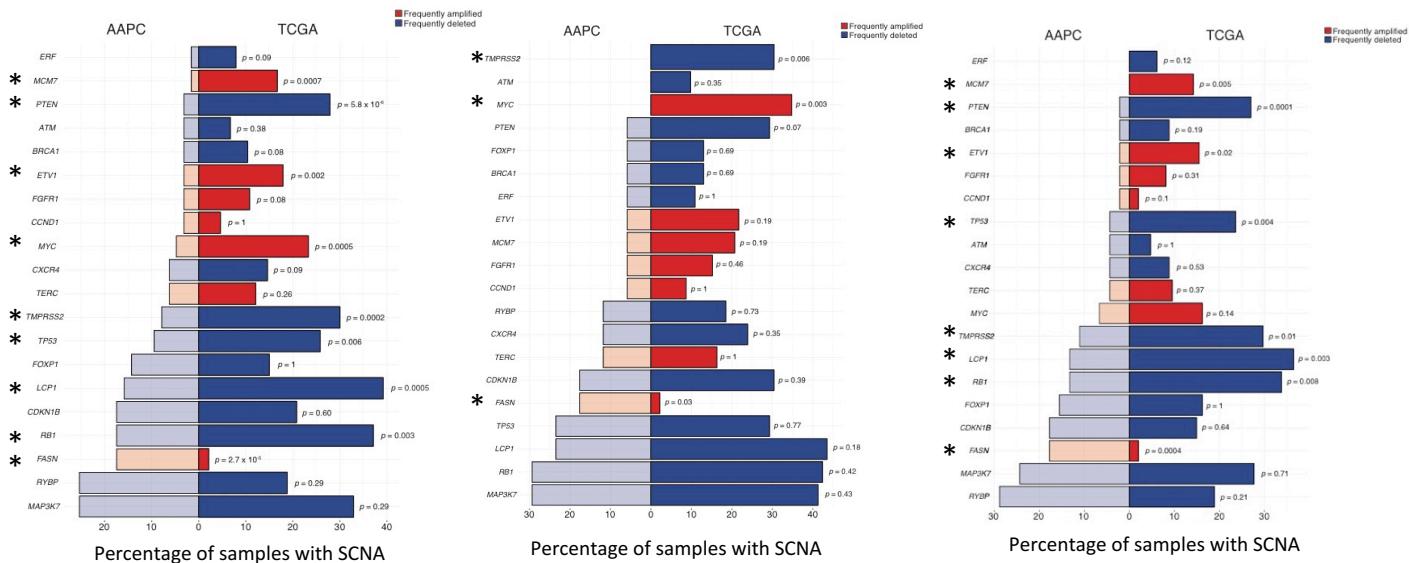


Supplemental Figure 2

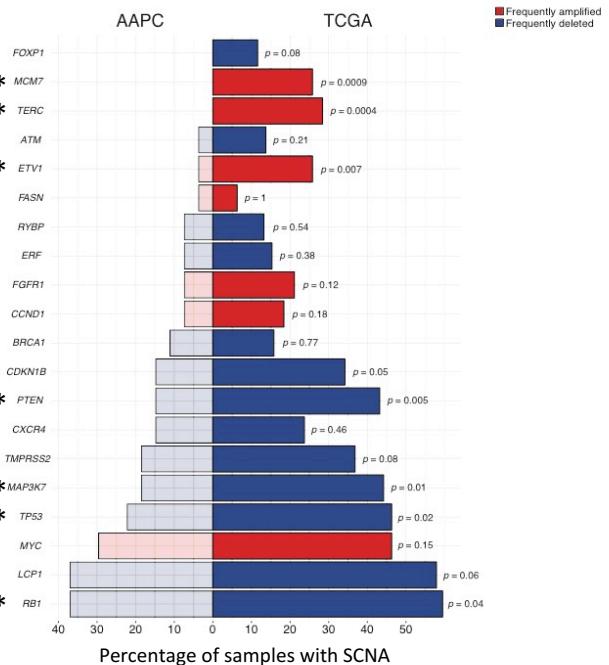
Lower
AAPC SCNA
threshold



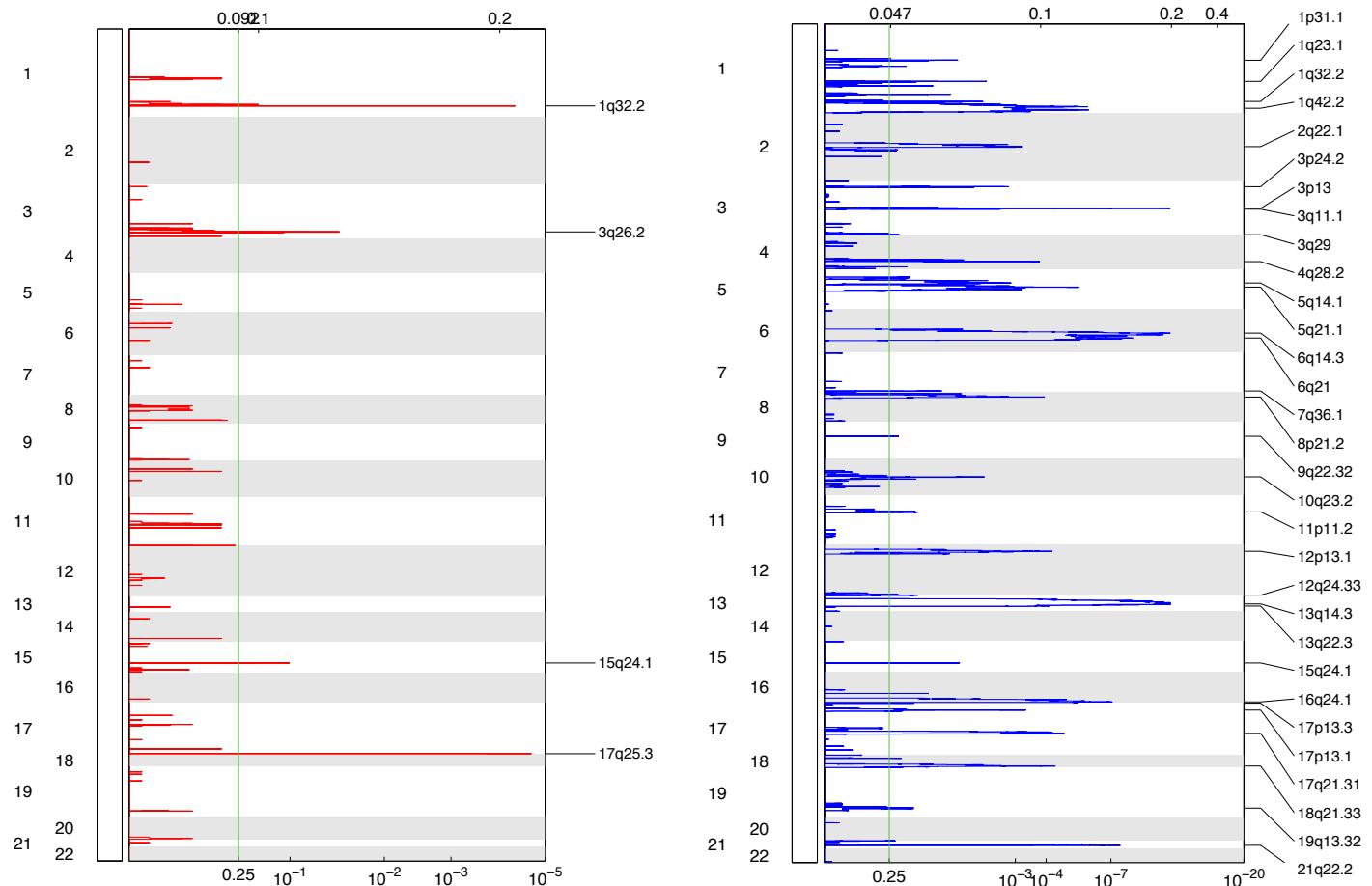
Gleason 7



Gleason 8+

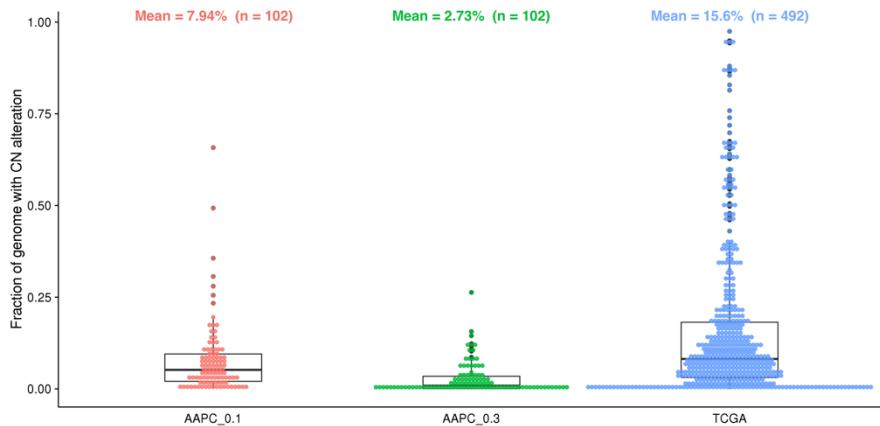


Supplemental Figure 3

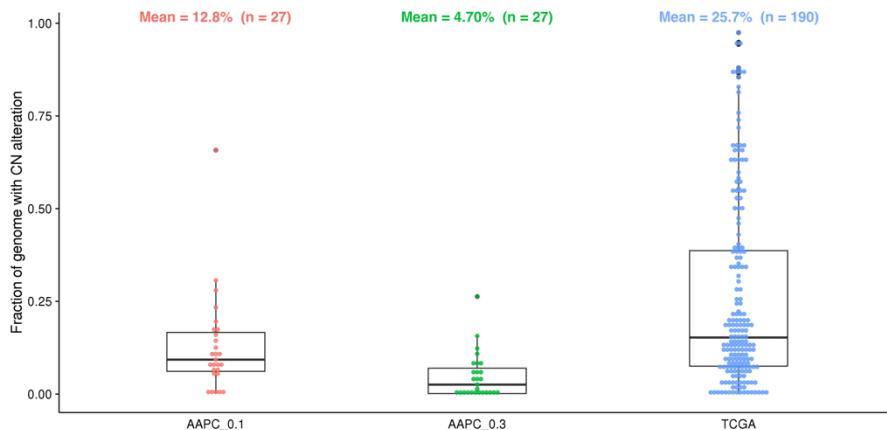


Supplemental Figure 4

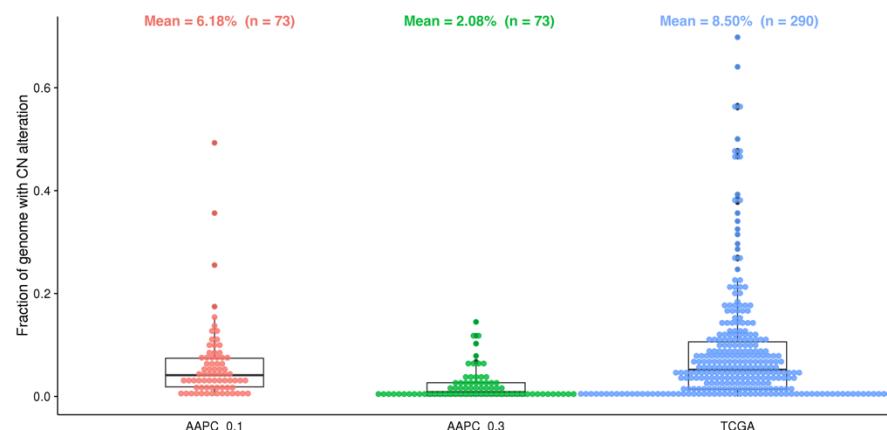
All tumors



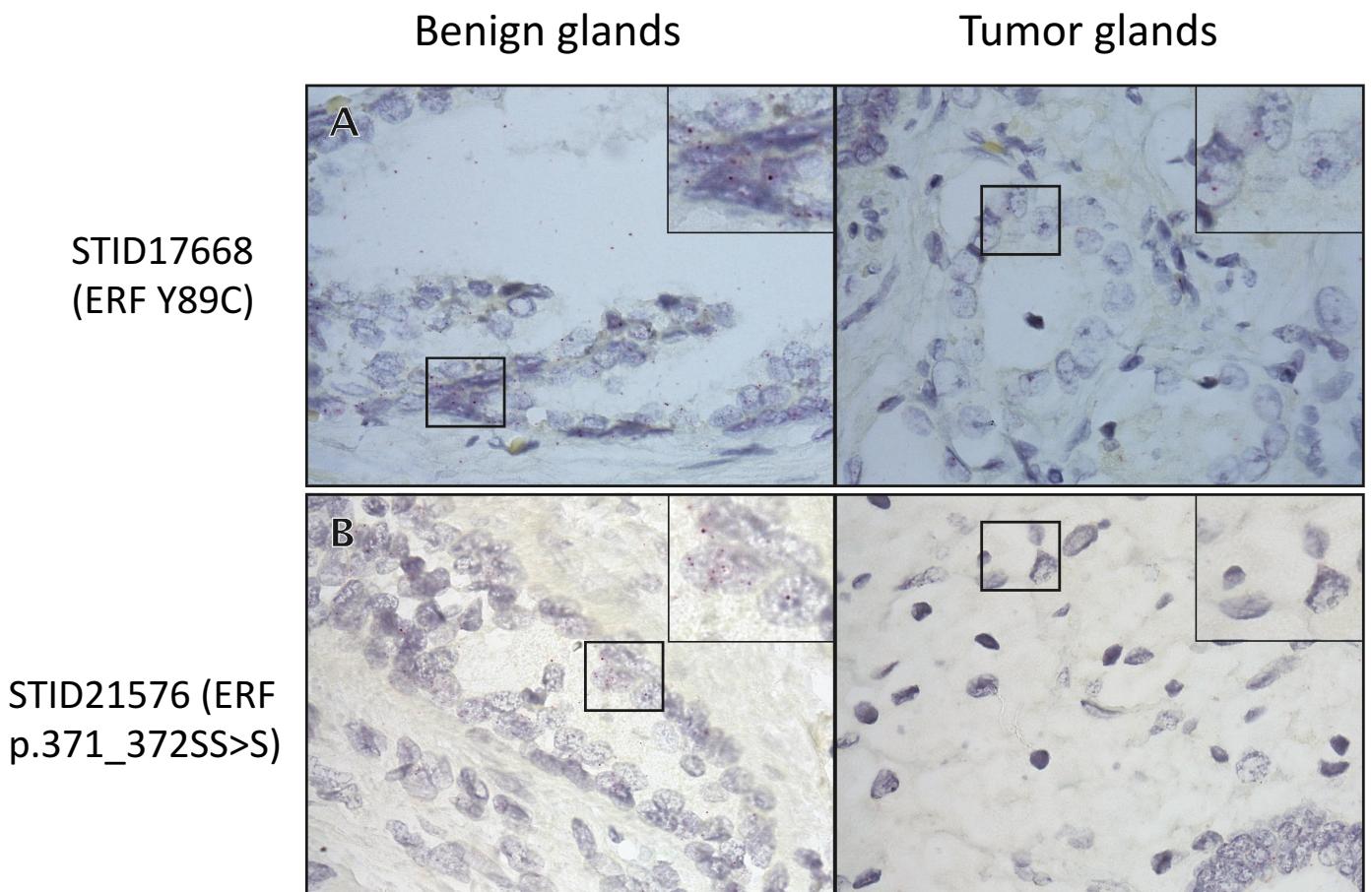
Gleason 8 and higher tumors



Gleason 7 and lower tumors

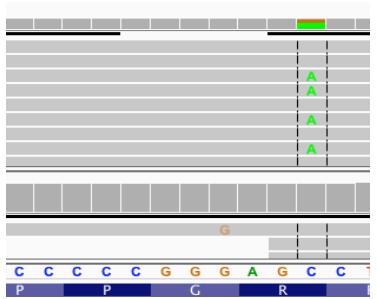


Supplemental Figure 5

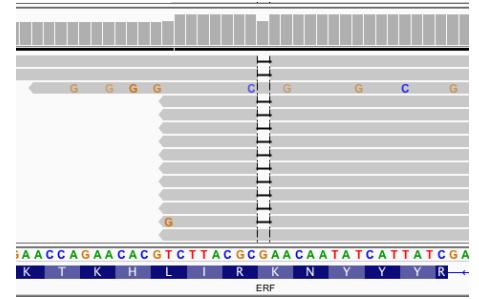


Supplemental Figure 6

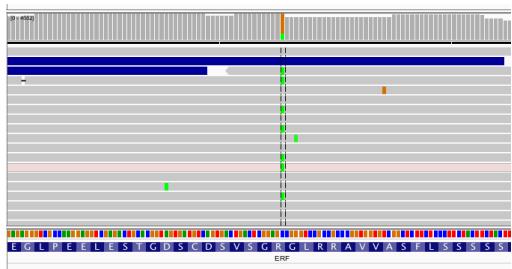
R218*



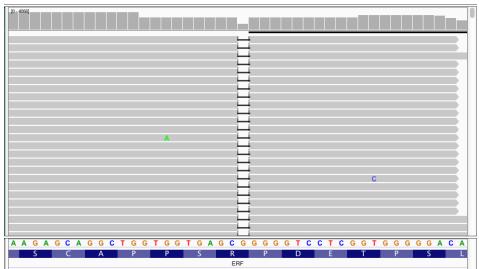
K91fs



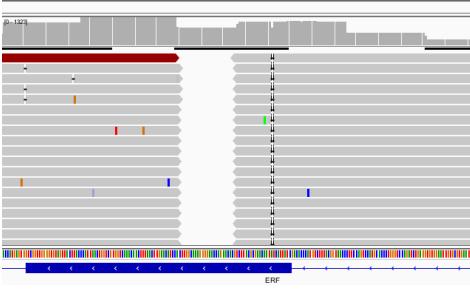
R183*



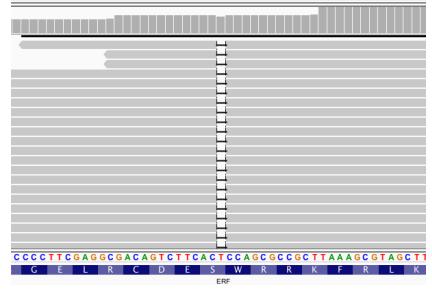
R160fs



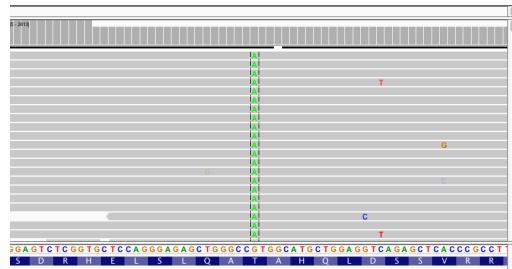
Y89fs



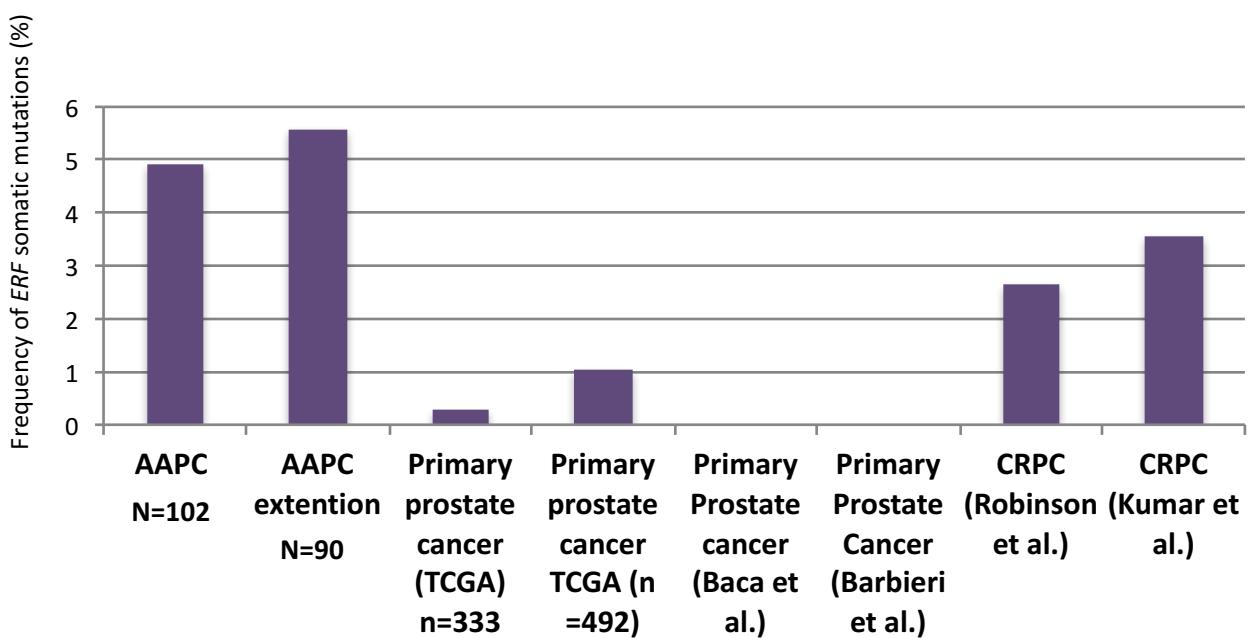
S489fs



T538M

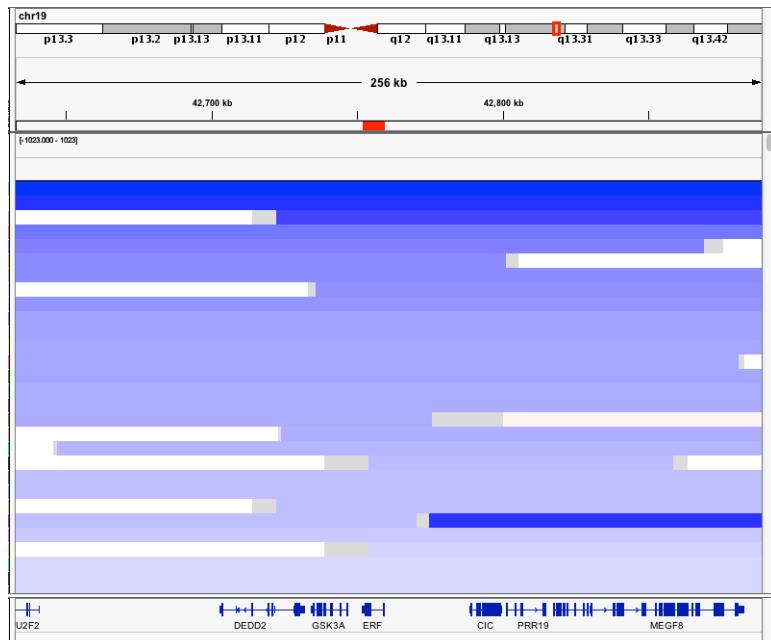


Supplemental Figure 7



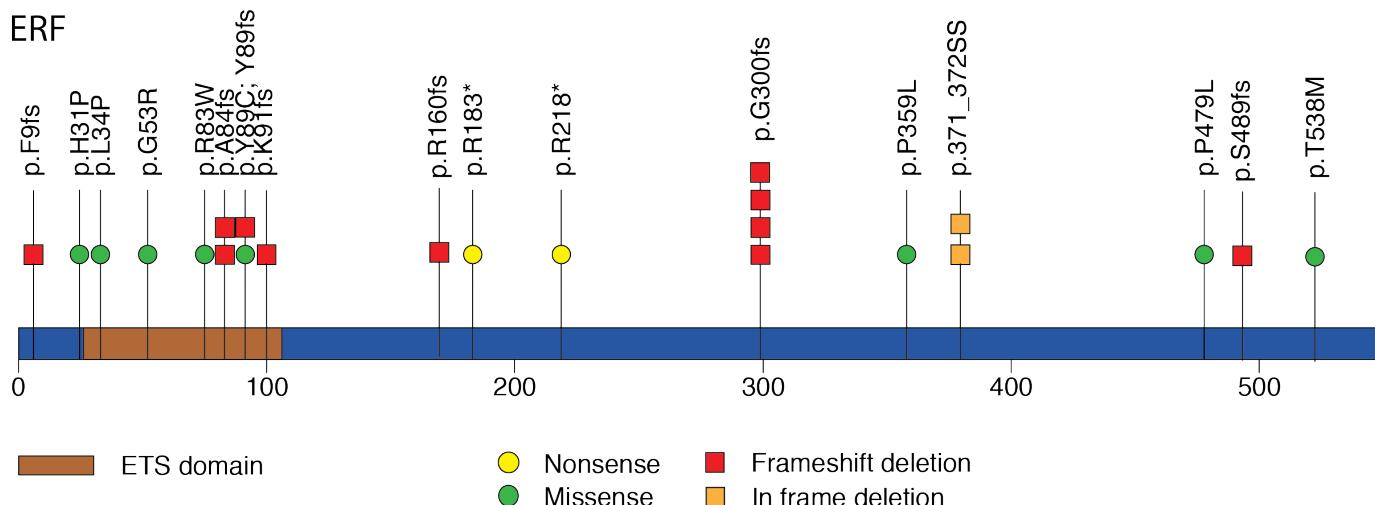
Supplemental Figure 8

A.

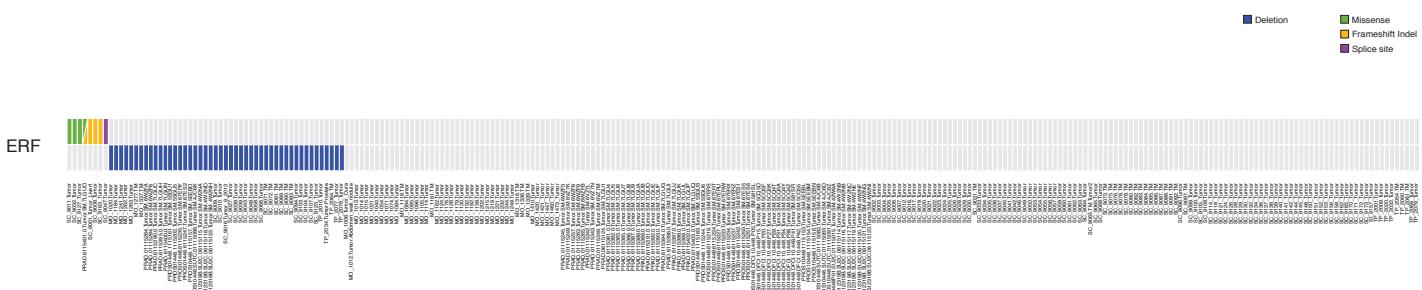


B.

ERF



C.



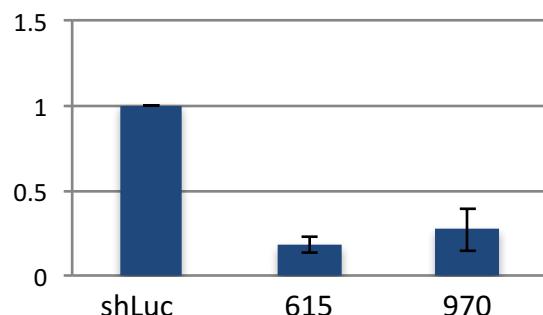
Supplemental Figure 9

Gleason Score:	6	7	8	9	10
ERF del=0: Observed Expected Ratio	43	217	55	100	2
	39.63	211.65	53.14	109.88	2.70
	1.085	1.025	1.035	0.910	0.741
ERF del=1: Observed Expected Ratio	1	18	4	22	1
	4.37	23.35	5.86	12.12	0.298
	0.228	0.771	0.683	1.82	3.36

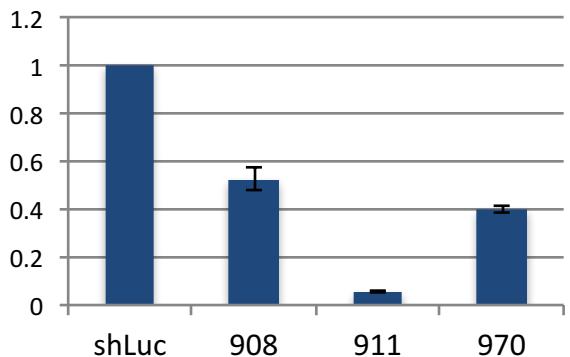
Supplemental Figure 10

Relative expression of ERF

PC-3

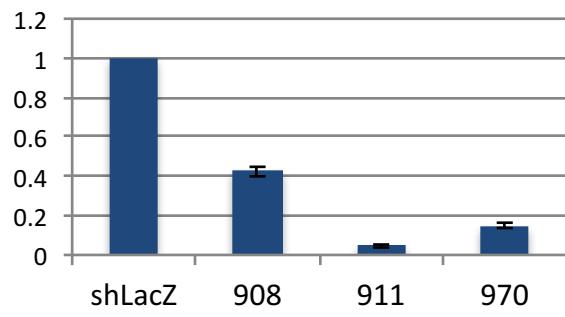


LHS-AR

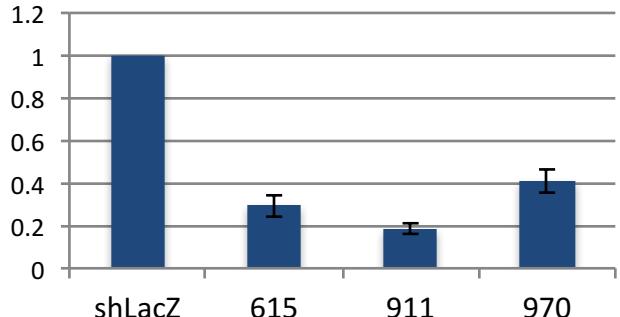


LNCaP

Relative expression of ERF

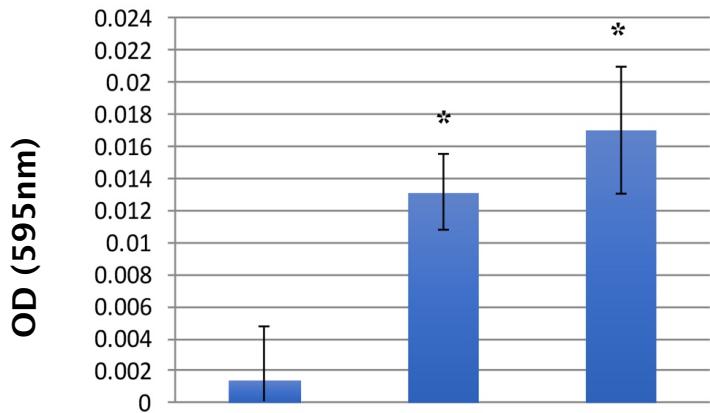
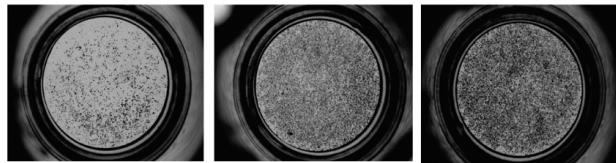


VCaP

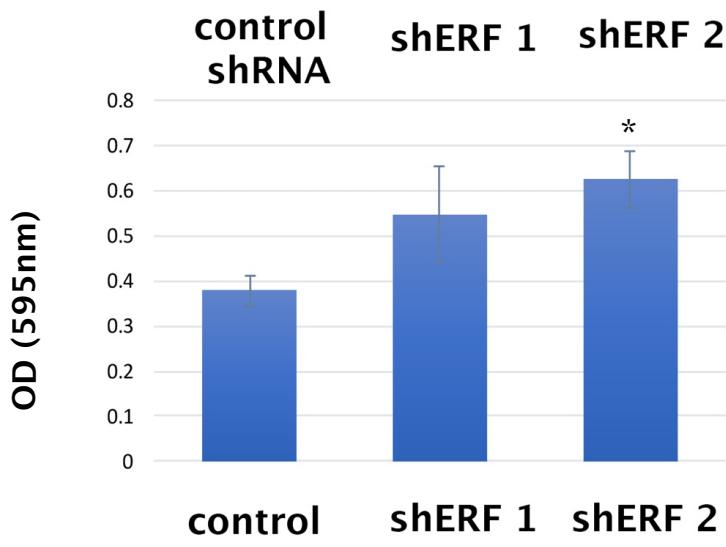


Supplemental Figure 11

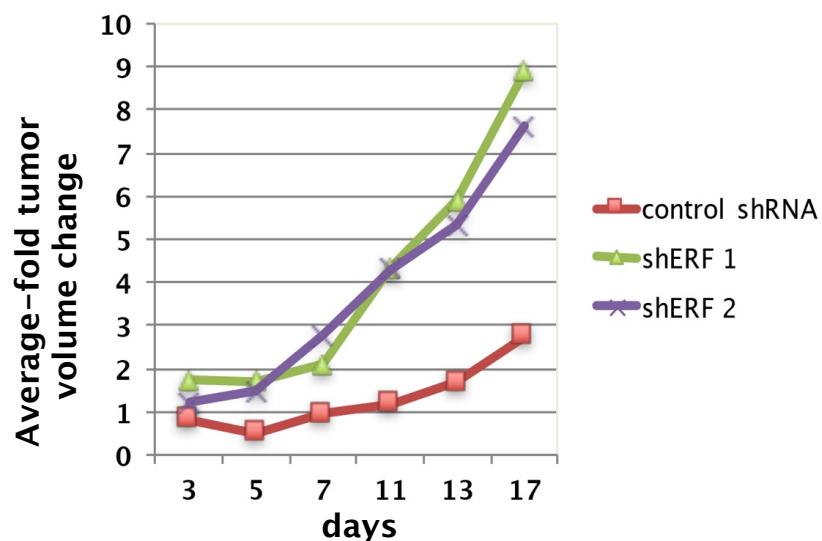
A.



B.

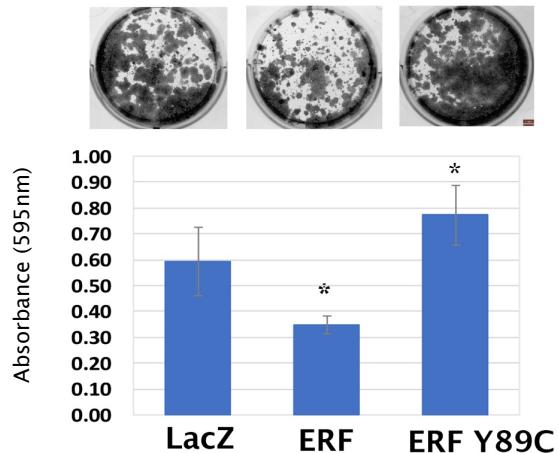


C.

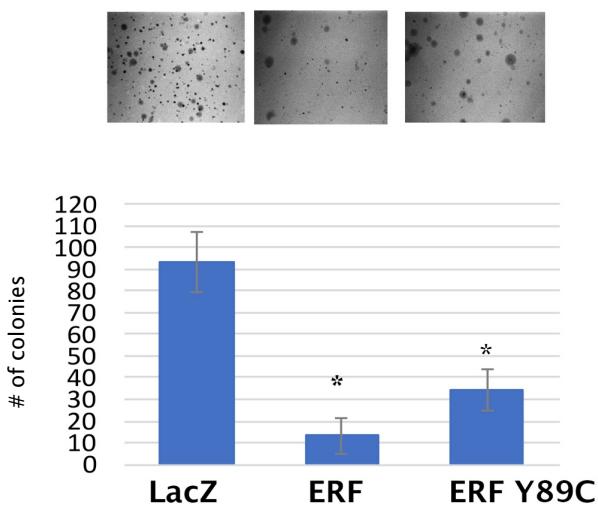


Supplemental Figure 12

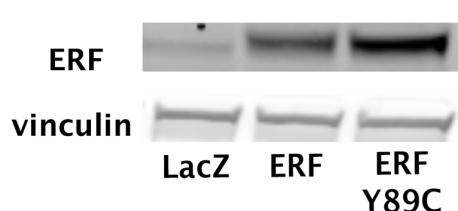
A.



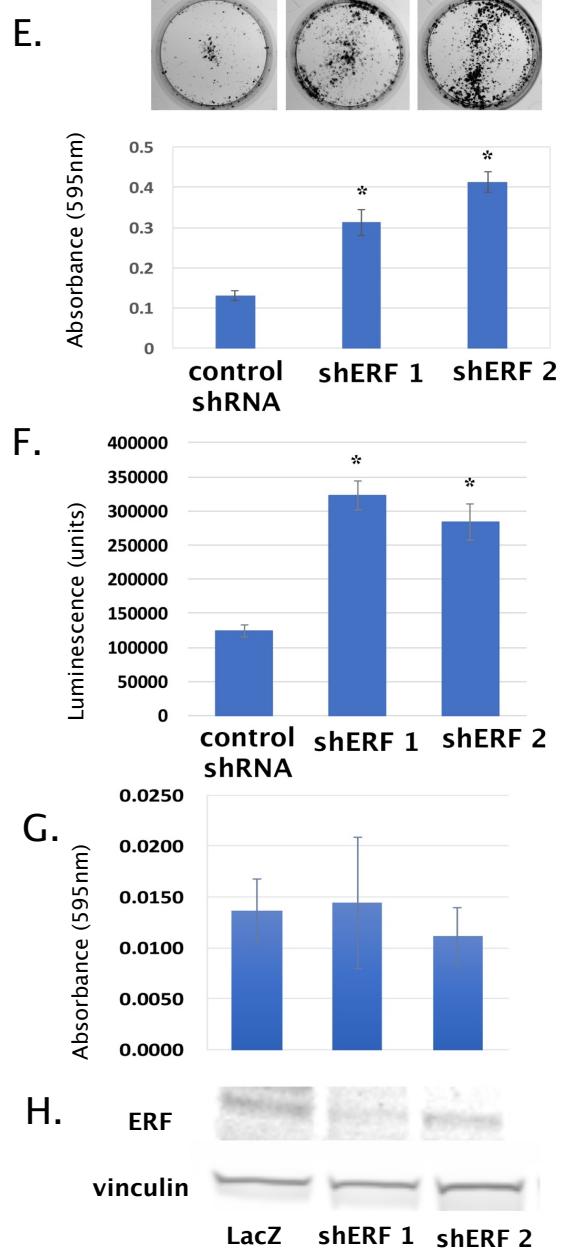
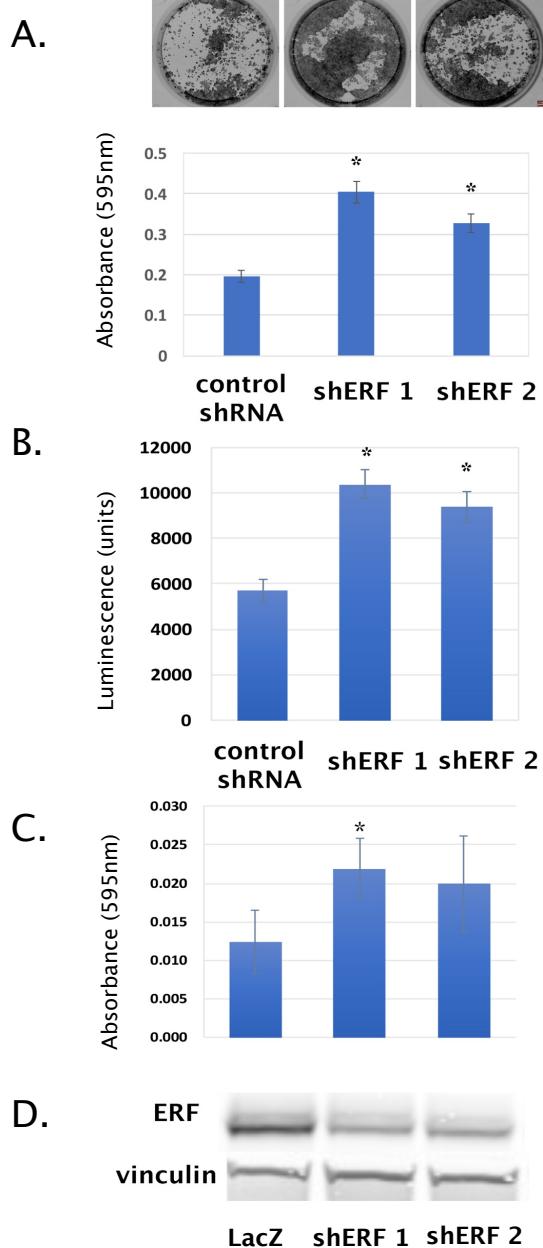
B.



C.

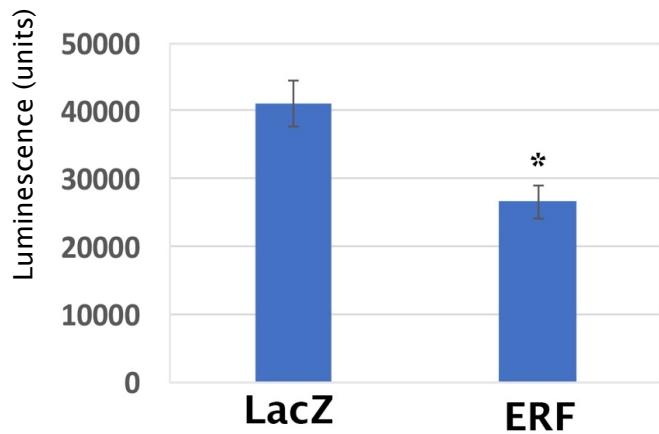
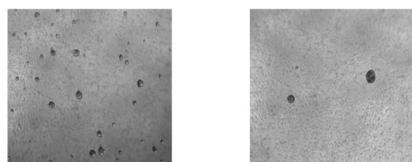


Supplemental Figure 13

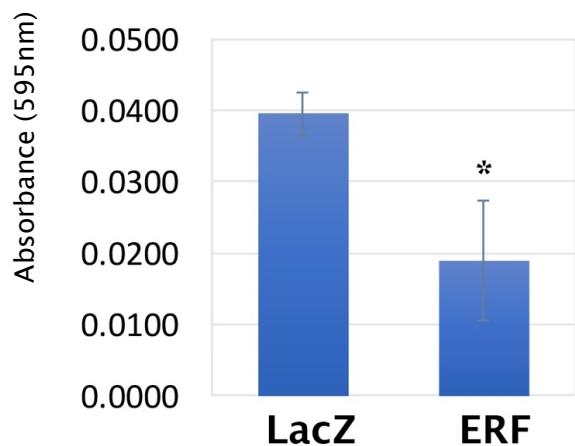
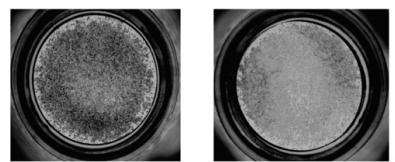


Supplemental Figure 14

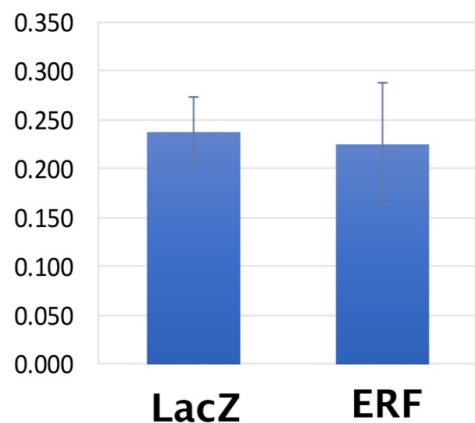
A.



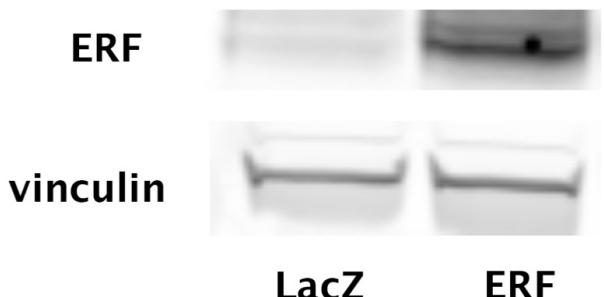
B.



C.



D.



Supplemental Figure 15

CCLE

ERG Pathway Expression

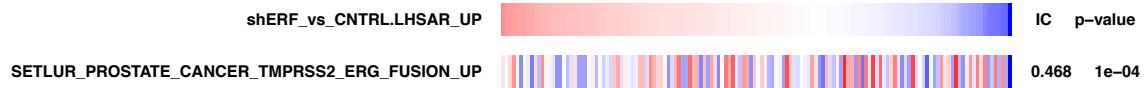


ETV1 Pathway Expression

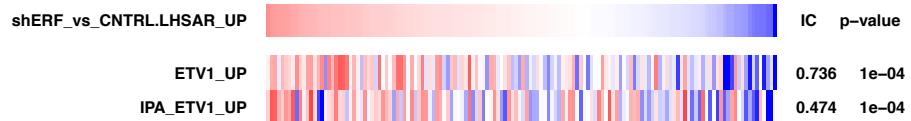


CRPC

ERG Pathway Expression



ETV1 Pathway Expression

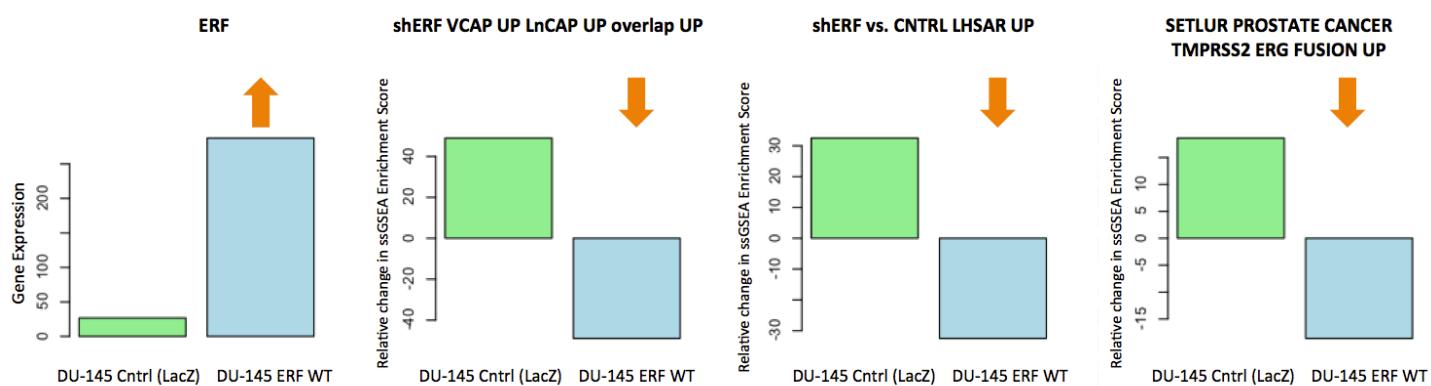


Supplemental Figure 16

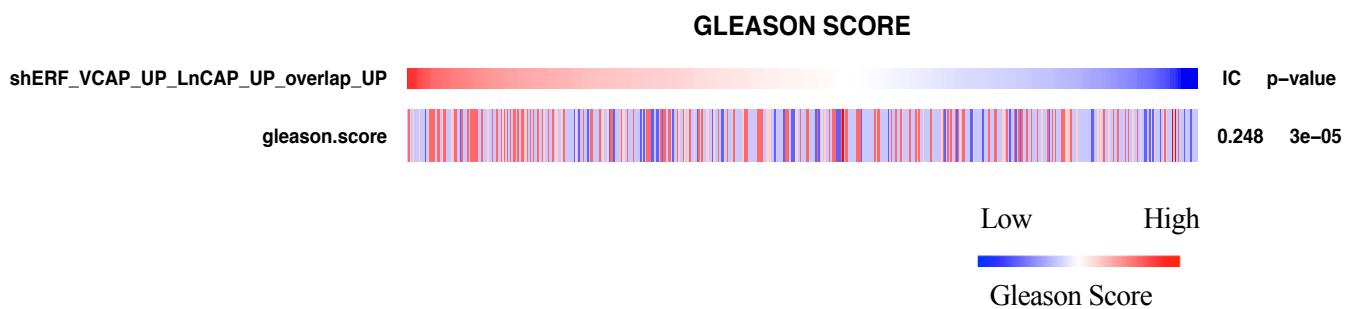


Supplemental Figure 17

A.



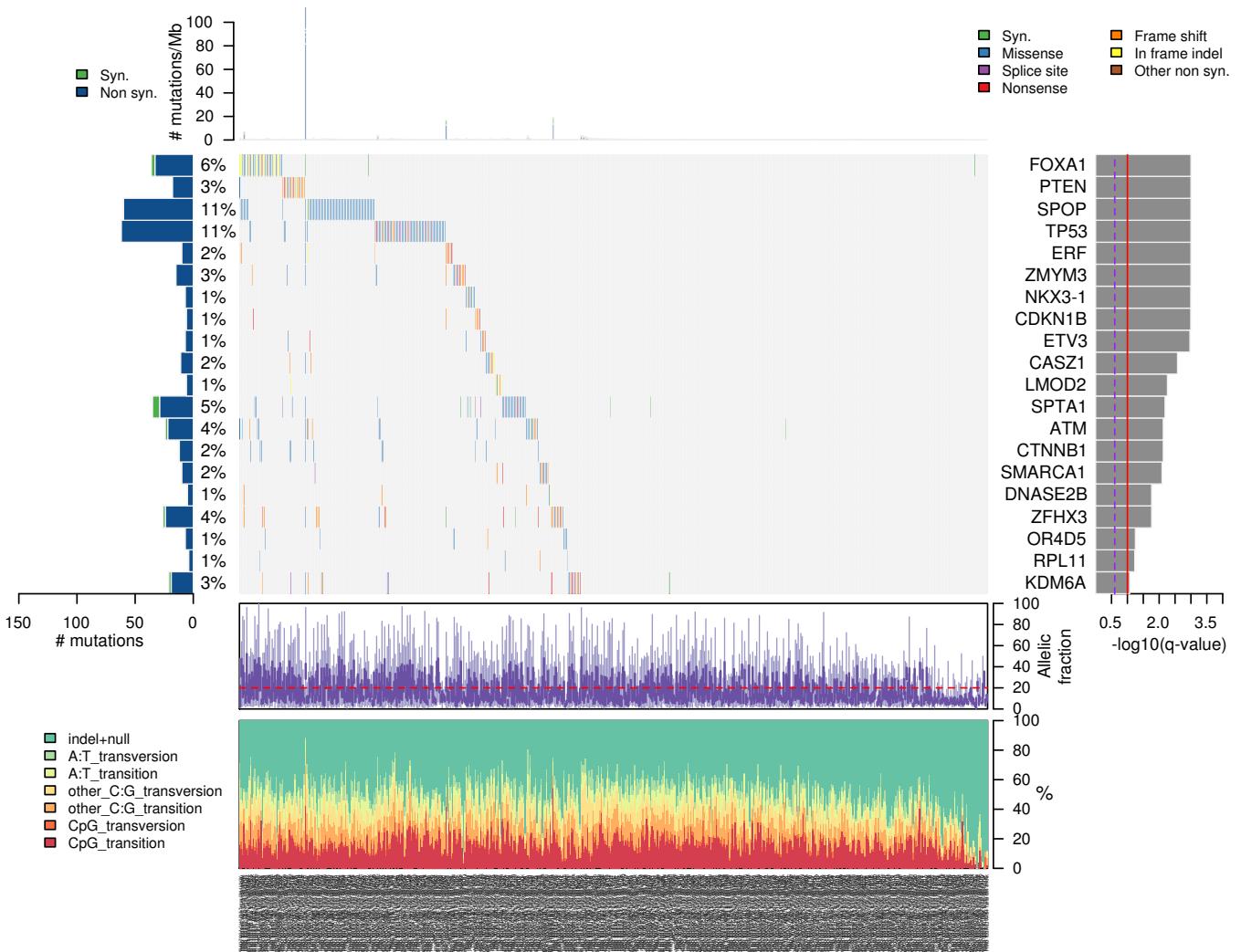
B.



Supplemental Figure 18



Supplemental Figure 19



Supplemental Figure 20

Place probesets on gene sequence

CE is BLUE , LE is RED and BL is green

>gi|666335577|ref|NM_006494| Homo sapiens Ets2 repressor factor (**ERF**), transcript variant 1,mRNA. [Homo sapiens]

gtgcgatattaacccgggaggcgccggcggggaggggagaggctctgagaggcgaggccgggtga
ggcggcgagggcgcccgacgggcgcggacgggacgggcagcgagggcgccggagccgcggc
ccggaatcggggcgcttcgccccggggcccccagcatgaagaccccggggacacagggttgcc
ttcccccattgggcctacaagccagactcgccccctggctcaaggcagatccagctgtggcactt
tatcctggagctgctgccaaggaggataccaggcgtcattgcctggcagggggactacgggg
aattcgtcatcaaagaccctgatgaggtggcccgctgtggggcggtcgcaagtcaagcccaag
atgaattacgacaagctgagccggccctgcgttattactataacaagcgcattctgcacaagac
caagggaaacgggtcacctacaagttcaatttcaacaaactgggtcggtcaattaccattca
ttgatgtgggggtggctgggggtgcagtgcggccagagtgcggccactgtccgtcggtggtagc
cacttccgcttccctccctcaacgcggccctccggaggtgtgtccccccaccggaggaccccgctcacc
accagcctgctttcatcttcatttcccttctccggctgtgtggcccccggctggccgag
gctcagtcagtgcactgttagtgcacgtcagactggagggaaaccgctggagggatcccgcc
ggccgaccaccggccctccggatctgggtgcctccggagggcccccgtggcccgctggccctggccca
tgaccctgggtcttcgagtcatacccgccctcggtggccctgaaccctcagcccttc
ctgtgtcgccctcgccggctcgatccctgctgcgtccctcaagtcctcccgcttc
acgcggccaccacccacccatcgccctacactccctcgccacgcgtgagcccgatgtaccccgatgggtggccg
ggggcccgccgcgtcaggggggaggctccacttctccctcagccctgaggacatgaaacgggtacc
tgcaggcccacacccaaacgcgtctacaactaccacctcagcccccgcgccttcgtcactaccct
gggctgggtggcccccagcccccacgcgcctgacaagtgcggctgcccgcgcgcgcgc
ccacccggccctccctccggctcgatccctttcttcctccctccattcaagttaa
tccagcccccactcgacgcggcagcggcagctggggagaaggccgtagccgtgt
aagagcggtgtggcagtgccaggccgtggctgagggggcaggggcgttagcccccacccggcc
accacagatcaagggtggagccatctcggaaggcgagtcggaggaggtagaggtgactgacatca
gtgatgaggatgaggaagacggggagggtgtcaagacgcggccgtgcaccccttaag
ccctgagcccgccgaggcaccggcatccctcgatccctcaagctacgcgtttaagcggcg
ctggagtgaagactgtcgccctcgaaagggtggggggccctggggctttgaggatgagggtg
aggacaagaagggtgcgtggggaggggccctggggagggtggggggccctcacc
agctctgacccctccagcatgccacgcggccagctccctggagcaccgagact
ccctggcaggggacctgtgtcccgaccccccattcgatccctttgtgcctta
ggagggtggggcagctcttgcctccctgcctccctccctccctccat
aaaacttaattttttttaaaaatgggtgggggtgggtgggtgc
ccctgtctgtgggttctaagctcgccaaagggtggtagggggaggggagg
gggggtcacccattctgggaatttataatttgcatttgc
ttttcttattacaatcgcttaggaagtaaggccctgtctccctccctgt
ccctgtacccaccccgctcgcccaactccctcgatccctccct
ctccaggggccatgagtgccctagggttcataccca
caattttataatgaacccaaaattccatgtgttgggggtggggggcgg
ccgcccattggccacaaatctacaagtgccgtcatccctccct
ggtccaaacccttcatccctcgatccctaggactggccat
gaagggggtgcctgaaaccaactggaaaggccctgtgc
gtggggactgagggtgggggaggtagagggggcc
gctggagaatgtattcaaaaactttggaccccttggatgc
gatgaaaaaaa